

Amendments to the Specification:

Between pages 71 and 72 of the application, on a new page, please insert the Sequence Listing submitted herewith.

On page 15, please replace the paragraph spanning lines 12-22 with the following paragraph:

Figure 12. Figure 12 depicts the amino acid alignment and phylogenetic analysis of vertebrate BCL2 proteins. (A) Alignment of BCL2 and BCL-xL proteins. Alignments shown are from zebrafish (z) BCL2 (SEQ ID NO:1) and BCL-xL (SEQ ID NO:5), Xenopus (x) BCL2 (SEQ ID NO:2) and BCL-xL (SEQ ID NO:6), chicken (c) BCL2 (SEQ ID NO:3) and BCL-xL (SEQ ID NO:7), and human (h) BCL2 (SEQ ID NO:4) and BCL-xL (SEQ ID NO:8). Amino acid residues conserved among both BCL2 and BCL-xL family members are indicated (#) while amino acid residues conserved among only BCL2 proteins are noted (*). Conserved BH1, BH2, BH3, and BH4 domains are denoted by a single line above the alignment. Dashes denote gaps introduced to maximize alignment. Sequence alignments were made using Megalign. (B) Diagram showing conserved domain homologies when zebrafish bcl-2 is compared to the human BCL2 protein. (C) Phylogenetic analysis of vertebrate BCL2 and BCL-xL proteins.

**U.S. Application Serial No. 10/659,705
Response to Notice to File Missing Parts dated Dec. 24, 2003 and
Second Preliminary Amendment dated Feb. 24, 2004**

Amendments to the Drawings:

In response to the Notice to File Missing Parts mailed December 24, 2003, please delete Figures 1-20 (15 sheets of informal drawings) and replace with Figures 1-20 (24 sheets of formal drawings) filed concurrently herewith.

Attachment: Twenty four (24) sheets of Formal Drawings (Figures 1-20).